#2

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RAW SEQUENCE LISTING   DATE: 01/16/2002
PATENT APPLICATION: US/10/024,944   TIME: 18:40:12

Input Set : A:\EXT-073.ST25.txt
Output Set: N:\CRF3\01162002\J024944.raw

```
 3 <110> APPLICANT: EXACT Science Corporation
 4        Boles, T. Christian
 5        Weir, Lawrence
 6        Stone, Benjamin
 8 <120> TITLE OF INVENTION: Detection of Non-Viral Organisms With SRP RNA
10 <130> FILE REFERENCE: EXT-073
12 <140> CURRENT APPLICATION NUMBER: US/10/024,944
13 <141> CURRENT FILING DATE: 2001-12-19
15 <150> PRIOR APPLICATION NUMBER: US 60/090,063
16 <151> PRIOR FILING DATE: 1998-06-19
18 <150> PRIOR APPLICATION NUMBER: US 08/971,845
19 <151> PRIOR FILING DATE: 1997-08-08
21 <160> NUMBER OF SEQ ID NOS: 27
23 <170> SOFTWARE: PatentIn version 3.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 21
27 <212> TYPE: RNA
28 <213> ORGANISM: Escherichia coli
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31 gucagguccg gaaggaagca g                                          21
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36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Complementary sequence of conserved E. coli 4.5S RNA region
41        nucleotide 44-65
43 <400> SEQUENCE: 2
44 gctgcttcct tccggacctg ac                                         22
47 <210> SEQ ID NO: 3
48 <211> LENGTH: 21
49 <212> TYPE: DNA
50 <213> ORGANISM: Artificial Sequence
52 <220> FEATURE:
53 <223> OTHER INFORMATION: Description of Artificial Sequence: shortened probe derived
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54        complementary sequence of E. coli 4.5S RNA region 44-65
56 <400> SEQUENCE: 3
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60 <210> SEQ ID NO: 4
61 <211> LENGTH: 12
62 <212> TYPE: DNA
63 <213> ORGANISM: Artificial Sequence
65 <220> FEATURE:
66 <223> OTHER INFORMATION: Description of Artificial Sequence:a shorter probe derived
from
67        complementary sequence of E. coli 4.5S RNA region 44-65
69 <400> SEQUENCE: 4
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ENTERED

70 gctgcttcct tc   12

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Input Set : **A:\EXT-073.ST25.txt**
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 75 <212> TYPE: DNA
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 78 <220> FEATURE:
 79 <223> OTHER INFORMATION: Description of Artificial Sequence: a short probe derived
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 82 <400> SEQUENCE: 5
 83 gctgcttcct tccg                                                    14
 86 <210> SEQ ID NO: 6
 87 <211> LENGTH: 14
 88 <212> TYPE: DNA
 89 <213> ORGANISM: Artificial Sequence
 91 <220> FEATURE:
 92 <223> OTHER INFORMATION: Description of Artificial Sequence: a short probe derived
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 93      complementary sequence of E. coli 4.5S RNA region 44-65
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 96 gacctgacct ggta                                                    14
 99 <210> SEQ ID NO: 7
100 <211> LENGTH: 41
101 <212> TYPE: DNA
102 <213> ORGANISM: Artificial Sequence
104 <220> FEATURE:
105 <223> OTHER INFORMATION: Description of Artificial Sequence: adaptor probe
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111 <210> SEQ ID NO: 8
112 <211> LENGTH: 41
113 <212> TYPE: DNA
114 <213> ORGANISM: Artificial Sequence
116 <220> FEATURE:
117 <223> OTHER INFORMATION: Description of Artificial Sequence: adaptor probe
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120 gctgcttcct tccggacctg acaaaaacga taaaccaacc a               41
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125 <212> TYPE: DNA
126 <213> ORGANISM: Artificial Sequence
128 <220> FEATURE:
129 <223> OTHER INFORMATION: Description of Artificial Sequence: adaptor probe
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132 ggcacacgcg tcatctgc                                                18
135 <210> SEQ ID NO: 10
136 <211> LENGTH: 29
137 <212> TYPE: RNA
138 <213> ORGANISM: Escherichia coli
140 <400> SEQUENCE: 10
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146 <212> TYPE: DNA
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149 <220> FEATURE:
150 <223> OTHER INFORMATION: Description of Artificial Sequence: complementary sequence
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151   conserved E. coli 4.5S RNA region nucleotides 36-65   30
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157 <210> SEQ ID NO: 12
158 <211> LENGTH: 29
159 <212> TYPE: DNA
160 <213> ORGANISM: Artificial Sequence
162 <220> FEATURE:
163 <223> OTHER INFORMATION: Description of Artificial Sequence: acrydite capture probe
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166 <400> SEQUENCE: 12
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170 <210> SEQ ID NO: 13
171 <211> LENGTH: 18
172 <212> TYPE: DNA
173 <213> ORGANISM: Artificial Sequence
175 <220> FEATURE:
176 <223> OTHER INFORMATION: Description of Artificial Sequence: fluorescent sandwich
probe 2F   18
178 <400> SEQUENCE: 13
179 ggcacacgcg tcatctgc
182 <210> SEQ ID NO: 14
183 <211> LENGTH: 12
184 <212> TYPE: DNA
185 <213> ORGANISM: Artificial Sequence
187 <220> FEATURE:
188 <223> OTHER INFORMATION: Description of Artificial Sequence:alkaline phosphatase
conjugated reporter probe RP-2
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192 <222> LOCATION: (1)..(1)   /
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196 <220> FEATURE:
W--> 197 <221> NAME/KEY: n
198 <222> LOCATION: (2)..(2)   /
199 <223> OTHER INFORMATION: cm
202 <220> FEATURE:
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204 <222> LOCATION: (3)..(3)   '
205 <223> OTHER INFORMATION: um
208 <220> FEATURE:
W--> 209 <221> NAME/KEY: n
210 <222> LOCATION: (4)..(4)   /
211 <223> OTHER INFORMATION: gm
214 <220> FEATURE:
W--> 215 <221> NAME/KEY: n

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        216 <222> LOCATION: (5)..(5)
        217 <223> OTHER INFORMATION: cm
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        223 <223> OTHER INFORMATION: um
        226 <220> FEATURE:
W--> 227 <221> NAME/KEY: n
        228 <222> LOCATION: (7)..(7)
        229 <223> OTHER INFORMATION: um
        232 <220> FEATURE:
W--> 233 <221> NAME/KEY: n
        234 <222> LOCATION: (8)..(8)
        235 <223> OTHER INFORMATION: cm
        238 <220> FEATURE:
W--> 239 <221> NAME/KEY: n
        240 <222> LOCATION: (9)..(9)
        241 <223> OTHER INFORMATION: cm
        244 <220> FEATURE:
W--> 245 <221> NAME/KEY: n
        246 <222> LOCATION: (10)..(10)
        247 <223> OTHER INFORMATION: gm
        250 <220> FEATURE:
W--> 251 <221> NAME/KEY: n
        252 <222> LOCATION: (11)..(11)
        253 <223> OTHER INFORMATION: um
        256 <220> FEATURE:
W--> 257 <221> NAME/KEY: n
        258 <222> LOCATION: (12)..(12)
        259 <223> OTHER INFORMATION: cm
        262 <400> SEQUENCE: 14
        263 nnnnnnnnnn nn
        266 <210> SEQ ID NO: 15
        267 <211> LENGTH: 19
        268 <212> TYPE: DNA
        269 <213> ORGANISM: Artificial Sequence
        271 <220> FEATURE:
        272 <223> OTHER INFORMATION: Description of Artificial Sequence: gel-immobilized
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        274 <220> FEATURE:
W--> 275 <221> NAME/KEY: n
        276 <222> LOCATION: (7)..(7)
        277 <223> OTHER INFORMATION: cm
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W--> 281 <221> NAME/KEY: n
        282 <222> LOCATION: (8)..(8)
        283 <223> OTHER INFORMATION: gm
        286 <220> FEATURE:
W--> 287 <221> NAME/KEY: n
        288 <222> LOCATION: (9)..(9)
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/024,944

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       292 <220> FEATURE:
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       294 <222> LOCATION: (10)..(10)
       295 <223> OTHER INFORMATION: 2'-O-methyl adenosine ribonucleotide
       298 <220> FEATURE:
W-->   299 <221> NAME/KEY: n
       300 <222> LOCATION: (11)..(11)
       301 <223> OTHER INFORMATION: cm
       304 <220> FEATURE:
W-->   305 <221> NAME/KEY: n
       306 <222> LOCATION: (12)..(12)
       307 <223> OTHER INFORMATION: cm
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W-->   311 <221> NAME/KEY: n
       312 <222> LOCATION: (13)..(13)
       313 <223> OTHER INFORMATION: um
       316 <220> FEATURE:
W-->   317 <221> NAME/KEY: n
       318 <222> LOCATION: (14)..(14)
       319 <223> OTHER INFORMATION: gm
       322 <220> FEATURE:
W-->   323 <221> NAME/KEY: n
       324 <222> LOCATION: (15)..(15)
       325 <223> OTHER INFORMATION: 2'-O-methyl adenosine ribonucleotide
       328 <220> FEATURE:
W-->   329 <221> NAME/KEY: n
       330 <222> LOCATION: (16)..(16)
       331 <223> OTHER INFORMATION: cm
       334 <220> FEATURE:
W-->   335 <221> NAME/KEY: n
       336 <222> LOCATION: (17)..(17)
       337 <223> OTHER INFORMATION: cm
       340 <220> FEATURE:
W-->   341 <221> NAME/KEY: n
       342 <222> LOCATION: (18)..(18)
       343 <223> OTHER INFORMATION: um
       346 <220> FEATURE:
W-->   347 <221> NAME/KEY: n
       348 <222> LOCATION: (19)..(19)
       349 <223> OTHER INFORMATION: gm
       352 <400> SEQUENCE: 15                                          19
W-->   353 tttttttnnnn nnnnnnnnn
       356 <210> SEQ ID NO: 16
       357 <211> LENGTH: 19
       358 <212> TYPE: DNA
       359 <213> ORGANISM: Artificial Sequence
       361 <220> FEATURE:
       362 <223> OTHER INFORMATION: Description of Artificial Sequence: gel-immobilized
acrydite-modified capture probe CP-2
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Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/024,944

DATE: 01/16/2002
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Input Set : A:\EXT-073.ST25.txt
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```
L:12  M:270 C: Current Application Number differs, Replaced Current Application Number
L:13  M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:191 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:197 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:203 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:209 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:215 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:221 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:227 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:233 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:239 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:245 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:251 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:257 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:263 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:275 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:281 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:287 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:293 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:299 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:305 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:311 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:317 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:323 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:329 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:335 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:341 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:347 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:365 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:371 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:377 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:383 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:389 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:395 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:401 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:407 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:413 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:419 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:425 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:431 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:437 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:443 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:455 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:461 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:467 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:473 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:479 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
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VERIFICATION SUMMARY
PATENT APPLICATION: US/10/024,944

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Input Set : A:\EXT-073.ST25.txt
Output Set: N:\CRF3\01162002\J024944.raw

L:485 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:491 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:497 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:503 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:509 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:515 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:521 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:533 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:623 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:713 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:881 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27

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